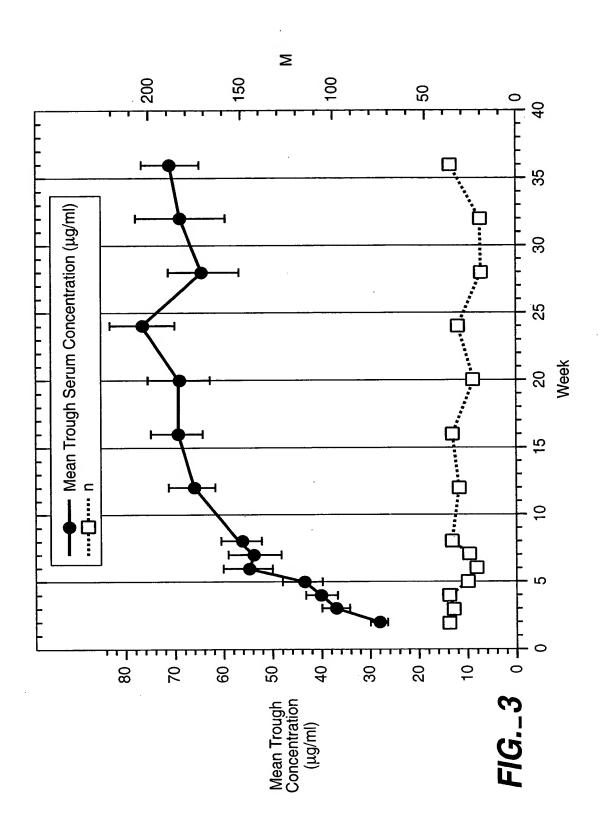
1/5 3H4 aa 541-599 4D5 aa 529-625 7C2 aa 22-53 7F3 aa 22-53 100 200 300 400 500 600 22-53 7C2 7F3 248 22 645 3H4 541 599 4D5 561 625 100 200 300 400 500 600 3H4 epitope (SEQ ID NO:8) 58 residues VEECRVLQGLPREYVNARHCLPCHPECQPQNGSVTCFGPEADQCVACAHYKDPPFCVAR 541 599 4D5 epitope (SEQ ID NO:9) 64 residues LPCHPECQPQNGSVTCFGPEADQCVACAHYKDPPFCVARCPSGVKPDLSYMPIWKFPDEEGACQP 561 625

## FIG.\_1

1	MELAALCRWGLLLALLPPGAA <b>STQVCTGTDMKLRLP</b>
38	SPETHLDMLRHLYQGCQVVQGNLELTYLPTNASLSFI
75	QDIQEVQGYVLIAHNQVRQVPLQRLRIVRGTQLFEDN
112	YALAVLDNGDPLNNTTPVTGASPGGLRELQLRSLTE
149	LKGGVLIQRNPQLCYQDTILWKDIFHKNNQLALTLII
186	TNRSRA

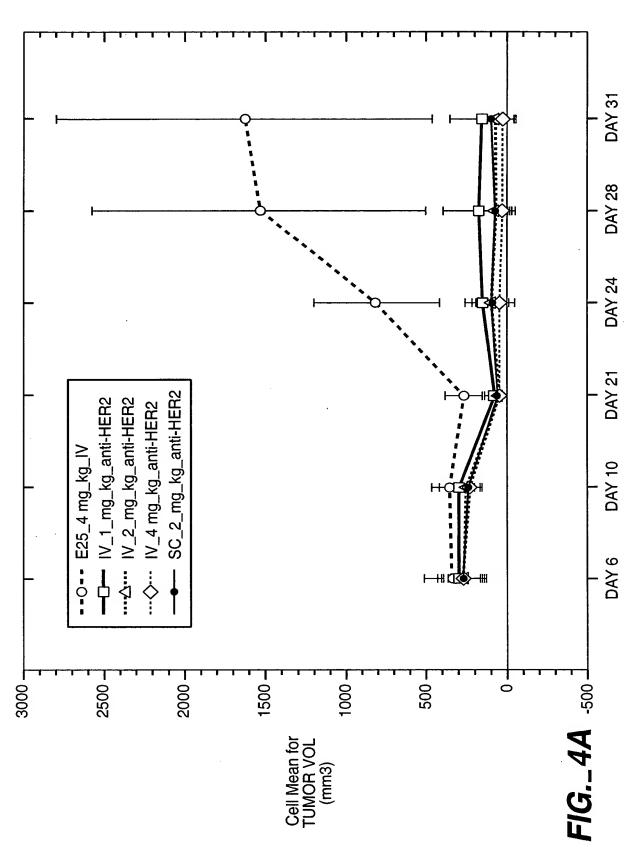
## FIG.\_2

2/5



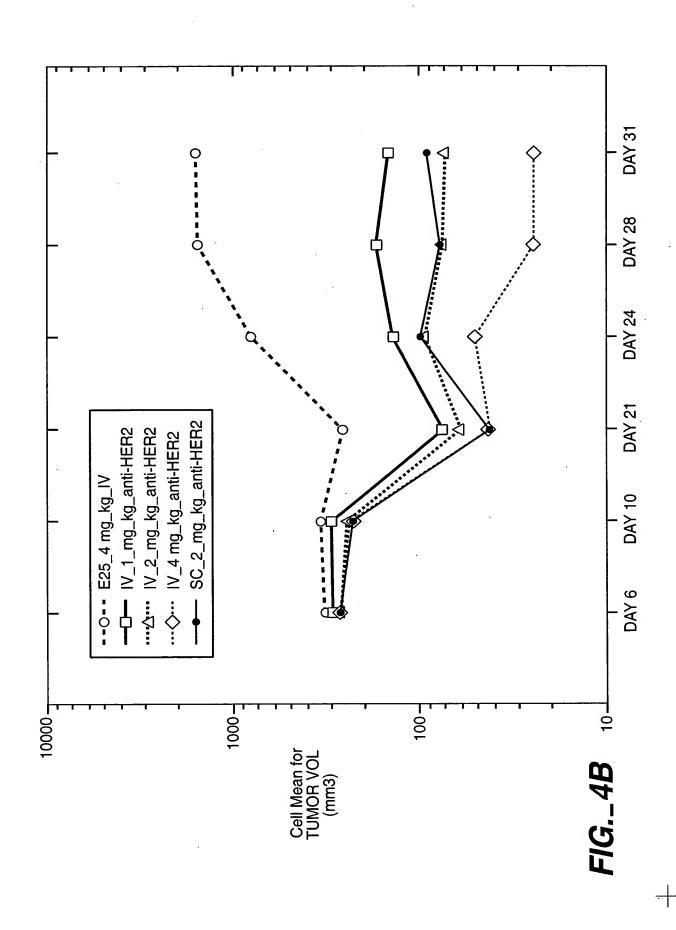
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3/5



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4/5



humIII

5/5

**VARIABLE LIGHT** 20 30 40 DTVMTQSHKIMSTSVGDRVSITC [KASQDVSIGVA----] WYQQRP .2C4 574 DIOMTOSPSSLSASVGDRVTITC [KASQDVSIGVA----] WYQQKP hum kI DIQMTQSPSSLSASVGDRVTITC [RASQSVSTSSYSYMH] WYQQKP 50 60 70 80 2C4 GOSPKLLIY [SASYRYT] GVPDRFTGSGSGTDFTFTISSVQA 574 GKAPKLLIY [SASYRYT] GVPSRFSGSGSGTDFTLTISSLQP GKAPKLLIY [AASSLES] GVPSRFSGSGSGTDFTLTISSLQP hum kI 100 90 2C4 EDLAVYYC [QQYYIYPYT] FGGGTKLEIK (SEQ ID NO:10) EDFATYYC [QQYYIYPYT] FGQGTKVEIK (SEQ ID NO:12) 574 EDFATYYC [QQYNSLPYT] FGQGTKVEIK (SEQ ID NO:14) hum kI FIG.\_5A VARIABLE HEAVY 10 20 30 40 2C4 EVOLOOSGPELVKPGTSVKISCKAS [GFTFTDYTMD] WVKQS 574 EVQLVESGGGLVQPGGSLRLSCAAS [GFTFTDYTMD] WVRQA EVOLVESGGGSVOPGGSLRLSCAAS [GFTFSSYAMS] WVRQA humIII 60 50 70 80 2C4 HGKSLEWIG [DVNPNSGGSIYNQRFKG] KASLTVDRSSRIVYM 574 PGKGLEWVA [DVNPNSGGSIYNQRFKG] RFTLSVDRSKNTLYL PGKGLEWVS [VISGDGGSTYYADSVKG] RFTISRDDSKNTLYL humIII 100 90 110 2C4 ELRSLTFEDTAVYYCAR [NLGPSFYFDY] WGQGTTLVTSS (SEQ ID NO:11) OMNSLRAEDTAVYYCAR [NLGPSFYFDY] WGOGTLVTVSS (SEO ID NO:13) 574

## FIG.\_5B

QMNSLRAEDTAVYYCAR [GRGGGS--DY] WGQGTLVTVSS (SEQ ID NO:15)